

Anuradha Wickramarachchi

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Scholar Uid: u-rFXwUAAAAJ

A passionate PhD candidate keen on learning state-of-the art and contributing towards the progress of humanity at large. Capable of teamwork and working under pressure.

WORK EXPERIENCE

November 2018 – present

Tutor

Tutoring Relational Database Systems.

School of Computing, Australian National University

December 2017 - July 2018

Software Engineer

HATCHYARD Pvt LTD, Sri Lanka

December 2017 - July 2018

Junior Consultant

Role of conducting research, student supervision and teaching.

Department of Computer Science and Engineering, University of Moratuwa, Sri Lanka

July 2016 - December 2016

Software Engineer (Intern)

CAKE LABS Pvt LTD, Sri Lanka

EDUCATION

August 2018 - present

PhD Candidate in Computer Science

School of Computing, Australian National University

Computational biology and bioinformatics

2014 - 2017

BSc (Hons) Computer Science and Engineering

Department of Computer Science and Engineering, University of Moratuwa, Sri Lanka

- GPA of 4.01/4.2
- · First class with deans list for all 8 semesters

TRAINING

Stanford Online

Machine Learning

An online non-credit course authorized by Stanford University and offered through Coursera

NVIDIA

Getting Started with AI on JETSON NANO

Offered through NVIDIA deep learning institute

DeepLearning.Al

Deep Learning Specialization (In-progress)

Neural networks and deep learning, Hyper-parameter tuning, Regularization and Optimization. Offered through Coursera

Udemy

PyTorch for Deep Learning with Python Bootcamp

Training on NumPy, Pandas and PyTorch including CNNs, LSTMs and GPU optimization.

PUBLICATIONS

First Author

- · LRBinner: Binning Long Reads in Metagenomics Datasets
 - Anuradha Wickramarachchi, Yu Lin (21st International Workshop on Algorithms in Bioinformatics (WABI) 2021) DOI: 10.4230/LIPIcs.WABI.2021.11
- PlasLR Enables Adaptation of Plasmid Prediction for Error-Prone Long Reads
 Anuradha Wickramarachchi, Vijini Mallawaarachchi, Lianrong Pu, Yu Lin (Under review at PLOS CB, Presented at GLBIO 2021) DOI: 10.1101/2021.06.14.448306
- GraphPlas: Refined Classification of Plasmid Sequences using Assembly Graphs Anuradha Wickramarachchi, Yu Lin (EEE/ACM Transactions on Computational Biology and Bioinformatics 2021) DOI: 10.1109/TCBB.2021.3082915
- MetaBCC-LR: Metagenomics Binning by Coverage and Composition for Long Reads Anuradha Wickramarachchi, Vijini Mallawaarachchi, Vaibhav Rajan, Yu Lin (Intelligent Systems for Molecular Biology (ISMB) 2020) DOI: 10.1093/bioinformatics/btaa441
- Use of nomadic computing devices for storage synchronization
 Anuradha Wickramarachchi, Dulaj Atapattu, Pamoda Wimalasiri, Ravidu Mallawa Arachchi, Gihan Dias (International Conference on Information Networking (ICOIN) 2018) DOI: 10.1109/ICOIN.2018.8343204

Collaborative Work

• Improving metagenomic binning results with overlapped bins using assembly graphs

Vijini G Mallawaarachchi, *Anuradha S Wickramarachchi*, Yu Lin (Algorithms Mol Biol 16, 3 (2021)) DOI: 10.1186/s13015-021-00185-6

 GraphBin2: Refined and Overlapped Binning of Metagenomic Contigs Using Assembly Graphs

Vijini G Mallawaarachchi, *Anuradha S Wickramarachchi*, Yu Lin (20th International Workshop on Algorithms in Bioinformatics; WABI 2020) DOI: 10.4230/LIPIcs.WABI.2020.8

- GraphBin: refined binning of metagenomic contigs using assembly graphs
 Vijini Mallawaarachchi, Anuradha Wickramarachchi, Yu Lin (Bioinformatics 2020) DOI: 10.1093/bioinformatics/btaa180
- Experiential Learning in Bioinformatics-Learner Support for Complex Workflow Modelling and Analysis.

Vijini Mallawaarachchi, *Anuradha Wickaramarachchi*, Anuradha Weliwita, Indika Perera, Dulani Meedeniya (International Journal of Emerging Technologies in Learning) DOI: 10.3991/ijet.v13i12.8608

Managing complex workflows in bioinformatics: An interactive toolkit with GPU acceleration

Anuradha Welivita, Indika Perera, Dulani Meedeniya, *Anuradha Wickramarach-chi*, Vijini Mallawaarachchi (IEEE transactions on nanobioscience) DOI: 10.1109/TNB.2018.2837122

SOFTWARE/TOOLS

Research Related

Freely available at github.com/anuradhawick

- LRBinner (PacBio/ONT reads clustering tool)
- MetaBCC-LR (PacBio/ONT reads clustering tool)
- PlasLR (Plasmid classification extension for PacBio/ONT reads)
- GraphPlas (Graph based plasmid classification extension tool)
- Seq2Vec fast memory mapped sequence vectoriser written in C++ (Boost)
- Seq2CovVec fast coverage vector computations for long reads written in C++ and Python

Software Engineering

Freely available at github.com/anuradhawick

- Vinyl.LK (AWS Serverless/Atlas Mongo hosted website for Sri Lankan vinyl records)
- YOLO Object Annotation Maker

AWARDS/ACHIEVEMENTS

Academic

- ANU Merit Fee Remission Scholarship
- Deans list for all 8 semesters during BSc
- Mahapola Sri Lankan undergraduate scholarship (merit based)
- GCE Advanced Level Island Rank 68 (District Rank 4)

Non-Academic

Awards

- Merit Award National Best Quality ICT Awards (NBQSA) Oct 2017 (Sri Lanka) for Final Year Honours Project
- Merit Award National Best Quality ICT Awards (NBQSA) Oct 2016 (Sri Lanka) for Taxi Master Project

Hackathons

- First place: IDEA Challenge 2017
- Google IO 2016 1st Runners up May 2016
- First place: IDEA Challenge 2016

SKILLS

Programming

Python, C, C++, JAVA, JavaScript, NodeJS, etc

- Industrial experience in ReactJS, MongoDB, AWS Serverless and AWS developer console.
- Research experience in python, C and C++.

Tools/Frameworks

Bioinformatics tools

- Aligners minimap(2), BLAST+, Edlib, etc
 Bioinformatics/Progamming libraries
- libminimap2, GATK, Edlib, biopython, etc
- Boost C++ library
 Machine Learning
- · Tensorflow, Pytorch, Sci-kit learn

Languages

English professional proficiency and Sinhalese native proficiency.

ADDITIONAL INFORMATION

Interests Blogging, Travelling and Gaming

Referees Available upon request.